MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HG <u>NKS</u> ARRKA	YNGSRRNIFS	12
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQ <u>NTS</u> HGSK	16
AQEIMKEAC K	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	20
HVPNQQDRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	24
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		27

Figure 1. Deduced amino acid sequence of Xenopus cerberus protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the Xenopus organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

	CAAGTCGCTC					60
CTTAAGGGTC	GTTCAGCGAG	TCTTTGTGAC	GTCCCAGATC	TATAGTATGT	TACAATGATT	
	GATCTGTATT					120
TACATGAGTC	CTAGACATAA	TAGCAGACGG	AACACTTACT	ACCTCGTCCT	TTTGTGAGTC	
	AAGGACAAAA					180
TTCCTGCTCT	TTCCTGTTTT	TGTATAAGTG	AATTGTCGTC	TCCAATGAAG	TCTTTTCTTT	
	TAGGAGCAAG					240
CTCCTCGTGC	ATCCTCGTTC	TAAGACGACC	ACTTATGATT	TCCAGAACTA	CTTGGGGTGT	
TTGGGCATGG	TGATTTTCGC	TTAGTAGCTG	AACTATTTGA	TTCCACCAGA	ACACATACAA	300
AACCCGTACC	ACTAAAAGCG	AATCATCGAC	TTGATAAACT	AAGGTGGTCT	TGTGTATGTT	
ACAGAAAAGA	GCCAGACATG	AACAAAGTCA	AGCTTTTCTC	AACAGTTGCC	CATGGAAACA	360
TGTCTTTTCT	CGGTCTGTAC	TTGTTTCAGT	TCGAAAAGAG	TTGTCAACGG	GTACCTTTGT	
AAAGTGCAAG	AAGAAAAGCT	TACAATGGTT	CTAGAAGGAA	TATTTTTCCT	CGCCGTTCTT	420
TTTCACGTTC	TTCTTTTCGA	ATGTTACCAA	GATCTTCCTT	ATAAAAAGGA	GCGGCAAGAA	
	AAATACAGAG					480
AACTATTTTC	TTTATGTCTC	CAATGACTTT	TCGGACCACG	GTTCTACAAG	ACCTTGTTAA	
TTTTGGTTAA	AATGAATGGA	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCGGT	ACCGTCATTT	CGTGTCCTTT	
Taatgaaaga	AGCTTGCAAA	ACCTTGTTTT	TCACTCAGAA	TATTGTACAT	GAAAACTGTG	600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	
	GATACAGAAC					660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTTAC	GTAGAGAGAG	GTACAAGGTT	
	TCGACGAAAT					720
TAGTCGTTCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
	GAATTGTACT					780
TGGACTGCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
					AACATGGATA	840
TTACGTGCAC	ACTTCGAGTA	TTCTCGTTGA	AGGTGGTTTG	ACGTGTCAAA	TTGTACCTAT	
					CTTTTGTTGG	900
GTAGATGATG	GGACGTGGTA	ATTTCCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	
					ATATAACCAC	960
•	•				TATATTGGTG	
					CATGCAAGAT	1020
TACCTTATTC	CTAACATACT	TAATATTAAT	TGTTTACCGT	AAAACACATI	GTACGTTCTA	

CTCTGTTCCA	TCAGTTGCAA	GATAAAAGGC	AATATTTGTT	TGACTTTTTT	TCTACAAAAT	1080
GAGACAAGGT	AGTCAACGTT	CTATTTTCCG	TTATAAACAA	ACTGAAAAAA	AGATGTTTTA	
GAATACCCAA	ATATATGATA	AGATAATGGG	GTCAAAACTG	TTAAGGGGTA	ATGTAATAAT	1140
CTTATGGGTT	TATATACTAT	TCTATTACCC	CAGTTTTGAC	AATTCCCCAT	TACATTATTA	
AGGGACTAAG	TTTGCCCAGG	AGCAGTGACC	CATAACAACC	AATCAGCAGG	TATGATTTAC	1200
TCCCTGATTC	AAACGGGTCC	TCGTCACTGG	GTATTGTTGG	TTAGTCGTCC	ATACTAAATG	
TGGTCACCTG	TTTAAAAGCA	AACATCTTAT	TGGTTGCTAT	GGGTTACTGC	TTCTGGGCAA	1260
ACCAGTGGAC	AAATTTTCGT	TTGTAGAATA	ACCAACGATA	CCCAATGACG	AAGACCCGTT	
AATGTGTGCC	TCATAGGGGG	GTTAGTGTGT	TGTGTACTGA	ATAAATTGTA	TTTATTTCAT	1320
TTACACACGG	AGTATCCCCC	CAATCACACA	ACACATGACT	TATTTAACAT	AAATAAAGTA	
TGTTACAAAA	AAAAAAA					
ACAATGTTTT	TTTTTTT					

Fig. 2. (Continuation page 2, SEQ ID NO:2).

MSRTRKVDSL	LLLAIPGLAL	LLLPNAYCAS	CEPVRIPMCK	SMPWNMTKMP	nhlhhstqan	60
AILAIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSVCER	ARAGCEPILI	120
KYRHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DFSMDSNNGN	CGSGREHCKC	180
KPMKATQKTY	LKNNYNYVIR	AKVKEVKVKC	HDATAIVEVK	EILKSSLVNI	PKDTVTLYTN	240
SGCLCPQLVA	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKKVKRWD	QKLRRPRKSK	300
DPVAPIPNKN	SNSRQARS					

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the Xenopus organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCCTT TCACACAGGA CTCCTGGCAG AGGTGAATGG TTAGCCCTAT G	GATTIGGTT 60
CTTAAGGGAA AGTGTGTCCT GAGGACCGTC TCCACTTACC AATCGGGATA CO	
TGTTGATTTT GACACATGAT TGATTGCTTT CAGATAGGAT TGAAGGACTT G	GATTTTAT 120
ACAACTAAAA CTGTGTACTA ACTAACGAAA GTCTATCCTA ACTTCCTGAA C	CTAAAAATA
CTAATTCTGC ACTITTAAAT TATCTGAGTA ATTGTTCATT TTGTATTGGA T	100 100
GATTAAGACG TGAAAATTTA ATAGACTCAT TAACAAGTAA AACATAACCT A	
OHISTONIO TOURNETTA AIRGOLIGIT INCONGIAN ANGAINCUL A	OCCIGATII
GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAACTA TAAGGTGGGG T	GAGTTGTAG 240
CTATTTGAAT TGAGGAACGA AAACTGAACG GGTATTTGAT ATTCCACCCC A	CTCAACATC
TTGCTTTTAC ATGTGCCCAG ATTTTCCCTG TATTCCCTGT ATTCCCTCTA A	
AACGAAAATG TACACGGGTC TAAAAGGGAC ATAAGGGACA TAAGGGAGAT T	TCATTCGGA
ACACATACAG GTTGGGCAGA ATAACAATGT CTCGAACAAG GAAAGTGGAC T	CATTACTGC 360
TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTC CTTTCACCTG A	
	.011211 01100
TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCCAA TGCTTACTGT G	CTTCGTGTG 420
ATGACCGGTA TGGACCTGAC CGCGAAGAGA ATAATGGGTT ACGAATGACA C	GAAGCACAC
AGCCTGTGCG GATCCCCATG TGCAAATCTA TGCCATGGAA CATGACCAAG A	
TCGGACACGC CTAGGGGTAC ACGTTTAGAT ACGGTACCTT GTACTGGTTC T	ACGGGTTGG
ATCTCCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT G	AAGGTTTGC 540
TAGAGGTGGT GTCGTGAGTT CGGTTACGGT AGGACCGTTA ACTTGTCAAA C	
TGACCACTGA ATGTAGCCAG GACCTTTTGT TCTTTCTGTG TGCCATGTAT G	
ACTGGTGACT TACATCGGTC CTGGAAAACA AGAAAGACAC ACGGTACATA C	CGGGGTAAA
GTACCATCGA TTTCCAGCAT GAACCAATTA AGCCTTGCAA GTCCGTGTGC G	
CATGGTAGCT AAAGGTCGTA CTTGGTTAAT TCGGAACGTT CAGGCACACG C	
william immediate criterium researcher chescarches c	.11100031
GGGCCGGCTG TGAGCCCATT CTCATAAAGT ACCGGCACAC TTGGCCAGAG A	AGCCTGGCAT 720
CCCGGCCGAC ACTCGGGTAA GAGTATTTCA TGGCCGTGTG AACCGGTCTC T	
GTGAAGAGCT GCCCGTATAT GACAGAGGAG TCTGCATCTC CCCAGAGGCT A	
CACTTCTCGA CGGCCATATA CTGTCTCCTC AGACGTAGAG GGGTCTCCGA T	ragcagtgtc
TGGAACAAGG AACAGATTCA ATGCCAGACT TCTCCATGGA TTCAAACAAT G	GAAATTGCG 840
ACCITGITCC TIGICIAAGI TACGGICTGA AGAGGIACCI AAGITTGITA C	
GAAGCGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGGC AACCCAAAAG A	
CTTCGCCGTC CCTCGTGACA TTTACGTTCG GGTACTTCCG TTGGGTTTTC T	TGCATAGAGT
	133mccco1cc
AGAATAATTA CAATTATGTA ATCAGAGCAA AAGTGAAAGA GGTGAAAGTG A TCTTATTAAT GTTAATACAT TAGTCTCGTT TTCACTTTCT CCACTTTCAC 1	
	111000100
ACGCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG A	AACATTCCTA 1020
TGCGTTGTCG TTAACACCTT CATTTCCTCT AAGAGTTCAG AAGGGATCAC I	

-						
	GACACTGTAC					1080
TTCTGTGTCA	CTGTGACATG	TGGTTGAGTC	CGACGAACAC	GGGGTCGAA	CAACGGTTAC	
AGGAATACAT	AATTATGGGC	TATGAAGACA	AAGAGCGTAC	CAGGCTTCTA	CTAGTGGAAG	1140
	TTAATACCCG			=		
	CGAAAAATGG		·			1200
CIAGGAACCG	GCTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
AGCTTCGACG	TCCCAGGAAA	AGCAAAGACC	CCGTGGCTCC	AATTCCCAAC	AAAAACAGCA	1260
TCGAAGCTGC	AGGGTCCTTT	TCGTTTCTGG	GGCACCGAGG	TTAAGGGTTG	TTTTTGTCGT	
**************************************	NCCCCCMN CM	macacmaacc	C11100m0m1	#001110#0#	1 m C C 1 C m m m C	1220
	AGCGCGTAGT					1320
TAAGGTCTGT	TCGCGCATCA	ATCTGATTGC	CTTTCCACAT	ACCTTTGAGA	TACCTGAAAC	
AAACTAAGAT	TTGCATTGTT	GGAAGAGCAA	AAAAGAAATT	GCACTACAGC	ACGTTATATT	1380
TTTGATTCTA	AACGTAACAA	CCTTCTCGTT	TTTTCTTTAA	CGTGATGTCG	TGCAATATAA	
	CTACAAGAAG	- -				1440
GATAACAAAT	GATGTTCTTC	GACCAAATCA	ACTAACATCA	AGAGGAAAGG	AAGAAAAAA	
TTATAACTAT	ATTTGCACGT	GTTCCCAGGC	AATTGTTTTA	TTCAACTTCC	AGTGACAGAG	1500
AATATTGATA	TAAACGTGCA	CAAGGGTCCG	TTAACAAAAT	AAGTTGAAGG	TCACTGTCTC	
	ATGTCTCAGC					1560
GTCACTGACT	TACAGAGTCG	GATTTCTTCG	AGTTAAGTAA	AGACTAGTTG	ATTACCACTG	
AAGTGTTTGA	TACTTGGGGA	AAGTGAACTA	ATTGCAATGG	TAAATCAGAG	AAAAGTTGAC	1620
	ATGAACCCCT					
CAATGTTGCT	TTTCCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
GTTACAACGA	AAAGGACATC	TACTTGTTCA	CTCTCTAGTG	TAAATTTACT	ACTAGTGAAA	
<i>ር</i> ርአ ጥጥጥ እ አጥ አ	CTTTCAGCAG	ጥጥጥአ ርጥጥአር	እም ር እር እጥር ጥ እ	CC3@CC3CC@	እ እ አ ጥ	1740
	GAAAGTCGTC					1/40
GGIAAAIIAI	GAAAGICGIC	AAAATCAATC	IACIGIACAI	CCIACGIGGA	IIIAGAIIIA	
ATTTTATCAT	AAATGAAGAG	CTGGTTTAGA	CTGTATGGTC	ACTGTTGGGA	AGGTAAATGC	1800
TAAAATAGTA	TTTACTTCTC	GACCAAATCT	GACATACCAG	TGACAACCCT	TCCATTTACG	
CTACTTTGTC	AATTCTGTTT	ፕ ልልልል ካጥር	Ставатават	ATTABGTCCT	***********	1860
	TTAAGACAAA					2000
AAAAAAAAA	AAAAA					
TTTTTTTTT	TTTTT					

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAIPM LLLGLMVLOT DCEIAOYYID EEEPPGTVIA VLSOHSIFNT TDIPATNFRL 60 MKQFNNSLIG VRESDGQLSI MERIDREQIC ROSLHCNLAL DVVSFSKGHF KLLNVKVEVR 120 DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT 180 RADGVKYADL VLMRELDREI OPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER 240 STIAVDLVED APLGYLLLEL HATDDDEGVN GEIVYGFSTL ASQEVRQLFK INSRTGSVTL 300 EGGVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY 360 IPETATKENF IALISTTDRA SGSNGOVRCT LYGHEHFKLQ QAYEDSYMIV TTSTLDRENI 420 AAYSLTVVAE DLGFPSLKTK KYYTVKVSDE NDNAPVFSKP QYEASILENN APGSYITTVI 480 ARDSDSDQNG KVNYRLVDAK VMGOSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA 540 DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD 600 EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA 660 TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FFVACTCKKK 720 AGEFKQVPEQ HGTCNEERLL STPSPQSVSS SLSQSESCQL SINTESENCS VSSNQEQHQQ 780 TGIKHSISVP SYHTSGWHLD NCAMSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK 840 RRALSSQCRH KPVLNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL

Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the Xenopus organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTCCCAG A			60
ACATTGCCAC ACTGTAACGGTG TO			120
AACTTTGATT C		 	 180
TGATGGTTTT A			240
CTGGCACTGT A GACCGTGACA T	 	 	 300
CAACCAATTT C			360
ATGGGCAGCT G TACCCGTCGA C			420
ACTGCAACCT G TGACGTTGGA C		 	 480
TGAAAGTGGA G ACTTTCACCT C		 	 540
ATGTGGAGGT G TACACCTCCA C			600
ATGAAGATGT T TACTTCTACA A			660
GCATTGATGT G		 	 720
AACTGGACAG G TTGACCTGTC C			780
TACCATCACT A ATGGTAGTGA T		 	 840
GCCCAGTGTT T			900
ACCTTTTGTT G			960
ATGGATTCAG C			1020

			CAAGCAGACT GTTCGTCTGA		1080
	· - · - · - ·	 	TACTTGTAAA ATGAACATTT		1140
· · · · · · · · · · · · · · · · · · ·		 	TACCCCTCTG ATGGGGAGAC		1200
			GAACTTTATA CTTGAAATAT		1260
			CTGTACTCTT GACATGAGAA		1320
		 	GATAGTTACC CTATCAATGG		1380
		 	TGCAGAAGAC ACGTCTTCTG		1440
		 	TGATGAGAAT ACTACTCTTA		1500
		 	AAATAATGCT TTTATTACGA		1560
		 	AAATGGCAAA TTTACCGTTT		1620
		 	ATTTGTTTCT TAAACAAAGA		1680
			AAAACTTAAA TTTTGAATTT		1740
		 	CACTCGCGTT GTGAGCGCAA	*	1800
		 	TAATCCTCTT ATTAGGAGAA		1860
				TTCCAGCTCA AAGGTCGAGT	1920
				CTGAGAGATC GACTCTCTAG	1980
		 		AAACAATTAA TTTGTTAATT	2040
				GGAAGACCTT CCTTCTGGAA	2100
				TCTAACGTTG AGATTGCAAC	2160

Fig. 6. (Continuation page 2, SEQ ID NO:6).

		GATCGATATG CTAGCTATAC		2220
		GGCCATCTTT CCGGTAGAAA		2280
	 	 TGAACAACAC ACTTGTTGTG		2340
		 CTCTTCTTCT GAGAAGAAGA		2400
• • • • • •		 TTGCAGCGTG AACGTCGCAC		2460
		TGTACCATCT ACATGGTAGA		2520
		ACATTCTCAC TGTAAGAGTG		2580
	 •	 AATGACAGTG TTACTGTCAC		2640
		CAGGCACAAG GTCCGTGTTC		2700
		TATTTCAGCC ATAAAGTCGG		2760
		AAGGGCTATA TTCCCGATAT		2820
		ATGCCTAACC TACGGATTGG	TGCACATACC ACGTGTATGG	2880
		CCTGTTGCTA GGACAACGAT	ATCGGATGCA TAGCCTACGT	2940
			AGAGATCGTC TCTCTAGCAG	3000
			ATCCTTCAGA TAGGAAGTCT	3060
			GCAAGTGCTT CGTTCACGAA	3120
			GGGGAGACAC CCCCTCTGTG	3180
			ATTTTTTGTT TAAAAAACAA	3240
• •	 		CTAACTAGCA GATTGATCGT	3300

Fig. 6. (Continuation page 3, SEQ ID NO:6).

	CAGACCTACA GTCTGGATGT					3360
	GGCCTTTTTA CCGGAAAAAT					3420
	GTCCTGAGTA CAGGACTCAT					3480
GTGATTTACA CACTAAATGT	CATAATAGGA GTATTATCCT	AACGCTTGGT TTGCGAACCA	TTCAGTGAAG AAGTCACTTC	TCTGTGTTGT AGACACAACA	ATATATTCTG TATATAAGAC	3540
	GCATTTTGTG CGTAAAACAC					3600
	TTGTAAATTA AACATTTAAT					

Fig. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60

TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120

PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC 180

KPVRATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240

SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300

TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

	•		•			
AAGCCTGGGA	CCATGGTCTG	CTGCGGCCCG	GGACGGATGC	TGCTAGGATG	GGCCGGGTTG	60
	GGTACCAGAC					
					1 CCCTCTC 1 C	120
	CTGCTCTCTG GACGAGAGAC					120
GATCAGGACC	GACGAGAGAC	GGACGAGGTC	CACGGGCCTC	GAGICCGACG	regorenere	
CCTGTCCGCA	TCCCGCTGTG	CAAGTCCCTT	CCCTGGAACA	TGACCAAGAT	GCCCAACCAC	180
	AGGGCGACAC					
					10000m00m0	240
	GCACCCAGGC					240
GACGTGGTGT	CGTGGGTCCG	ATTGCGGTAG	GACCGGTACC	TIGICAAGCI	ICCCGACGAC	
GGCACCCACT	GCAGCCCGGA	TCTTCTCTTC	TTCCTCTGTG	CAATGTACGC	ACCCATTTGC	300
	CGTCGGGCCT					
						2.60
	TCCAGCACGA					360
TGGTAGCTGA	AGGTCGTGCT	CGGGTAGTTC	GGGACGTTCA	GACACACACT	CGCGCGGGCT	
CAGGGCTGCG	AGCCCATTCT	CATCAAGTAC	CGCCACTCGT	GGCCGGAAAG	CTTGGCCTGC	420
	TCGGGTAAGA					
						400
GACGAGCTGC	CGGTGTACGA	CCGCGGCGTG	TGCATCTCTC	CTGAGGCCAT	CGTCACCGCG	480
CTGCTCGACG	GCCACATGCT	GGCGCCGCAC	ACGTAGAGAG	GACTCCGGTA	GCAGTGGCGC	
GACGGAGCGG	ATTTTCCTAT	GGATTCAAGT	ACTGGACACT	GCAGAGGGGC	AAGCAGCGAA	540
	TAAAAGGATA					
						600
	GTAAGCCTGT					600
GCAACGTTTA	CATTCGGACA	GTCTCGATGT	GICTICIGGA	TAAAGGCCTT	GITARIGITG	
TATGTCATCC	GGGCTAAAGT	TAAAGAGGTA	AAGATGAAAT	GTCATGATGT	GACCGCCGTT	660
	CCCGATTTCA					
						700
	AGGAAATTCT					720
CACCTTCACT	TCCTTTAAGA	TTTCCGTAGT	GACCATTIGI	AAGGTTCCCT	GIGGCAGIIA	
CTTTATACC	CCTCTGGCTG	CCTCTGTCCT	CCACTTACTG	TCAATGAGGA	ATATGTCATC	7 80
GAAATATGGT	GGAGACCGAC	GGAGACAGGA	GGTGAATGAC	AGTTACTCCT	TATACAGTAG	
						040
ATGGGCTAT	AAGACGAGGA	ACGTTCCAGG	TTACTCTTGG	TAGAAGGCTC	TATAGCTGAG	840
TACCCGATA	Trergereer	TGCAAGGTCC	. AATGAGAACC	AICIICCGAC	ATATCGACTC	
AAGTGGAAG	S ATCGGCTTG	TAAGAAAGTO	AAGCGCTGG	ATATGAAACT	CCGACACCTT	900
TTCACCTTC	C TAGCCGAAC	ATTCTTTCAC	TTCGCGACC	C TATACTTTG!	A GGCTGTGGAA	
					- CACCAACTCT	960
GGACTGGGT.	A AAACTGATG	TAGCGATTC	C ACTCAGAAT	C AGAAGTCTG(G CAGGAACTCT C GTCCTTGAGA	900
CCTGACCCA	T TITGACTAC	ATCGCTAAG	3 IGNGICITA	G ICIICNONCI	GICCIIONON	

;

						1000
AATCCCCGGC	CAGCACGCAG	CTAAATCCTG	AAATGTAAAA	GGCCACACCC	ACGGACTCCC	1020
TTAGGGGCCG	GTCGTGCGTC	GATTTAGGAC	TTTACATTTT	CCGGTGTGGG	TGCCTGAGGG	
					macma) 0003	1080
TTCTAAGACT	GGCGCTGGTG	GACTAACAAA	GGAAAACCGC	ACAGTTGTGC	TCGTGACCGA	1080
AAGATTCTGA	CCGCGACCAC	CTGATTGTTT	CCTTTTGGCG	TGTCAACACG	AGCACTGGCT	
			* * COMM COMMC	accmoccomm	mcmccmccmm	1140
TTGTTTACCG	CAGACACCGC	GTGGCTACCG	AAGTTACTTC	CCCACCCCAA	ACACCACCAA	1140
AACAAATGGC	GTCTGTGGCG	CACCGATGGC	TTCAATGAAG	GCCAGGGAA	AADDAEDADA	
	TGGGGTTAGA	ጥርርጥጥጥል ልጥል	ጥርጥጥልጥልጥልጥ	TCTGTTTCAT	CAATCACGTG	1200
CTTAATGGCG	ACCCCAATCT	ACCADATTAT	ACAATATATA	AGACAAAGTA	GTTAGTGCAC	
GAATTACCGC	ACCCCAATCI	AGGMM111111				
CCC A CጥርጥጥC	TTTTGCAACC	AGAATAGTAA	ATTAAATATG	TTGATGCTAA	GGTTTCTGTA	1260
CCCTGACAAG	AAAACGTTGG	TCTTATCATT	TAATTTATAC	AACTACGATT	CCAAAGACAT	
000101101210						
CTGGACTCCC	TGGGTTTAAT	TTGGTGTTCT	GTACCCTGAT	TGAGAATGCA	ATGTTTCATG	1320
GACCTGAGGG	ACCCAAATTA	AACCACAAGA	CATGGGACTA	ACTCTTACGT	TACAAAGTAC	
TAAAGAGAGA	ATCCTGGTCA	TATCTCAAGA	. ACTAGATATT	GCTGTAAGAC	AGCCTCTGCT	1380
ATTTCTCTCT	TAGGACCAGT	ATAGAGTTCT	TGATCTATAA	CGACATTCTG	TCGGAGACGA	
					. mamas s s amm	1440
GCTGCGCTTA	TAGTCTTGTG	TTTGTATGCC	TTTGTCCATI	TCCCTCATGC	TGTGAAAGTT	1440
CGACGCGAAT	ATCAGAACAC	AAACATACGG	AAACAGGTAA	AGGGAGTACG	ACACTTTCAA	
			, massamasa		GCAGAGTAGC	1500
ATACATGTTT	' ATAAAGGTAG	AACGGCATTT	TORRATURGE	CACIGCACAP	CGTCTCATCG	1500
TATGTACAAA	TATTTCCATC	TIGCCGIAA	ACITIAGIC	GIGACGIGIA		
002202002	·	ጥር አርር አ አ አ ር	CCACACAGC	TGACTTATT	TCAAGATTGG	1560
CCMACACCAC	TOTAL CALLES	1 ACTOCTTTG(GGTGTGTCG	ACTGAATAA	A AGTTCTAACC	
GGIIGIGGIC	. CIICGIAMI	nercerre				
CAGGCAGCA	AATAAATAGI	GTTGGGAGC	C AAGAAAAGA	A TATTTTGCC	r ggttaaggg	1620
GTCCGTCGT	TTATTTATCA	CAACCCTCG	G TTCTTTTCT	r ATAAAACGG	A CCAATTCCCC	
CACACTGGA	A TCAGTAGCCC	TTGAGCCAT	T AACAGCAGT	G TTCTTCTGG	C AAGTTTTTGA	1680
GTGTGACCT	r AGTCATCGG	AACTCGGTA	A TTGTCGTCA	C AAGAAGACC	G TTCAAAAACT	
						1540
TTTGTTCAT	A AATGTATTC	A CGAGCATTA	G AGATGAACT	T ATAACTAGA	C ATCTGTTGTT	1740
AAACAAGTA'	r ttacataag:	r gctcgtaat	C TCTACTTGA	A TATTGATCT	G TAGACAACAA	
			3 3 3 CCC3 PPPC	m m~~xm~~~~	C CTCTCC & TTCC	1800
ATCTCTATA	G CTCTGCTTC	TTCTAAATC	A AACCCATIG	A ACCUACCAC	C CTCTCCATTC G GAGAGGTAAG	1000
TAGAGATAT	C GAGACGAAG	S AAGATTTAG	1 TIGGGIAAC	A ACCIACGAG	- C1101100111110	

	TTGGCTTGCT					1860
TATTTATTTA	AACCGAACGA	CATAACCGGT	CCTTTTCTTT	CATAATTTCA	TACGTACGTA	
GTGCACCAGG	GTGTTATTTA	ACAGAGGTAT	GTAACTCTAT	AAAAGACTAT	AATTTACAGG	1920
CACGTGGTCC	CACAATAAAT	TGTCTCCATA	CATTGAGATA	TTTTCTGATA	TTAAATGTCC	
ACACGGAAAT	GTGCACATTT	GTTTACTTTT	TTTCTTCCTT	TTGCTTTGGG	CTTGTGATTT	1980
	CACGTGTAAA					
TGGTTTTTGG	TGTGTTTATG	TCTGTATTTT	GGGGGGTGGG	TAGGTTTAAG	CCATTGCACA	2040
	ACACAAATAC					
TTCAAGTTGA	ACTAGATTAG	AGTAGACTAG	GCTCATTGGC	CTAGACATTA	TGATTTGAAT	2100
	TGATCTAATC					
TTGTGTTGTT	TAATGCTCCA	TCAAGATGTC	TAATAAAAGG	AATATGGTTG	TCAACAGAGA	2160
	ATTACGAGGT					
CGACAACAAC	AACAAA					
 СС ФСФФСФТС	TTGTTT					

MVCGSPGGML	LLRAGLLALA	ALCLLRVPGA	RAAACEPVRI	PLCKSLPWNM	TKMPNHLHHS	60
TQANAILAIE	QFEGLLGTHC	SPDLLFFLCA	MYAPICTIDF	QHEPIKPCKS	VCERARQGCE	120
PILIKYRHSW	PENLACEELP	VYDRGVCISP	EAIVTADGAD	FPMDSSNGNC	RGASSERCKC	180
KPIRATQKTY	FRNNYNYVIR	AKVKEIKTKC	HDVTAVVEVK	EILKSSLVNI	PRDTVNLYTS	240
SGCLCPPLNV	NEEYIIMGYE	DEERSRLLLV	EGSIAEKWKD	RLGKKVKRWD	MKLRHLGLSK	300
SDSSNSDSTQ	SQKSGRNSNP	RQARN.				

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10. This sequence was assembled from public ESTs from the Genbank database (accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG CCGCCTCGCC	GCCTTTTGGC (CGGAAAACCG	GTCCACTGCG (CAGGTGACGC (CGGCTGCACC (GCCGACGTGG (CTGCCCCATC GACGGGGTAG	TGCCGGGATC ACGGCCCTAG	60
ATGGTCTGCG	GCAGCCCGGG .	AGGGATGCTG	CTGCTGCGGG	CCGGGCTGCT	TGCCCTGGCT	120
TACCAGACGC	CGTCGGGCCC	TCCCTACGAC	GACGACGCCC	GGCCCGACGA	ACGGGACCGA	
GCTCTCTGCC	TGCTCCGGGT	GCCCGGGGCT	CGGGCTGCAG	CCTGTGAGCC	CGTCCGCATC	180
CGAGAGACGG	ACGAGGCCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGCGTAG	
CCCCTGTGCA	AGTCCCTGCC	CTGGAACATG	ACTAAGATGC	CCAACCACCT	GCACCACAGC	240
GGGGACACGT	TCAGGGACGG	GACCTTGTAC	TGATTCTACG	GGTTGGTGGA	CGTGGTGTCG	
ACTCAGGCCA	ACGCCATCCT	GGCCATCGAG	CAGTTCGAAG	GTCTGCTGGG	CACCCACTGC	300
TGAGTCCGGT	TGCGGTAGGA	CCGGTAGCTC	GTCAAGCTTC	CAGACGACCC	GTGGGTGACG	
AGCCCCGATC	TGCTCTTCTT	CCTCTGTGCC	ATGTACGCGC	CCATCTGCAC	CATTGACTTC	360
TCGGGGCTAG	ACGAGAAGAA	GGAGACACGG	TACATGCGCG	GGTAGACGTG	GTAACTGAAG	
CAGCACGAGC	CCATCAAGCC	CTGTAAGTCT	GTGTGCGAGC	GGGCCCGGCA	GGGCTGTGAG	420
GTCGTGCTCG	GGTAGTTCGG	GACATTCAGA	CACACGCTCG	CCCGGGCCGT	CCCGACACTC	
CCCATACTCA	TCAAGTACCG	CCACTCGTGG	CCGGAGAACC	TGGCCTGCGA	GGAGCTGCCA	480
GGGTATGAGT	AGTTCATGGC	GGTGAGCACC	GGCCTCTTGG	ACCGGACGCT	CCTCGACGGT	
GTGTACGACA CACATGCTGT	GGGGCGTGTG	CATCTCTCCC GTAGAGAGGG	GAGGCCATCG CTCCGGTAGC	TTACTGCGGA AATGACGCCT	CGGAGCTGAT GCCTCGACTA	540
TTTCCTATGG	ATTCTAGTAA	CGGAAACTGT	AGAGGGGCAA	GCAGTGAACG	CTGTAAATGT	600
AAAGGATACG	TAAGATCATT	GCCTTTGACA	TCTCCCCGTT	CGTCACTTGC	GACATTTACA	
AAGCCTATTA	GAGCTACACA	GAAGACCTAT	TTCCGGAACA	ATTACAACTA	TGTCATTCGG	660
TTCGGATAA	CTCGATGTGT	CTTCTGGATA	AAGGCCTTGT	TAATGTTGAT	ACAGTAAGCC	
GCTAAAGTTA	A AAGAGATAAA	GACTAAGTGC	CATGATGTGA	CTGCAGTAGT	GGAGGTGAAG	720
CGATTTCAA	T TTCTCTATTT	CTGATTCACG	GTACTACACT	GACGTCATCA	CCTCCACTTC	
GAGATTCTA CTCTAAGAT	A AGTCCTCTCI T TCAGGAGAGA	GGTAAACATT CCATTTGTAA	CCACGGGACA	CTGTCAACCT GACAGTTGG	CTATACCAGC GATATGGTCG	780
TCTGGCTGC AGACCGACG	C TCTGCCCTCC G AGACGGGAGC	C ACTTAATGTT	AATGAGGAAT	TATATCATCAS A TATAGTAGTA	GGGCTATGAA CCCGATACTT	840

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	GTTCCAGATT					900
CTACTCCTTG	CAAGGTCTAA	TGAGAACCAC	CTTCCGAGAT	ATCGACTCTT	CACCTTCCTA	
CGACTCGGTA	AAAAAGTTAA	GCGCTGGGAT	ATGAAGCTTC	GTCATCTTGG	ACTCAGTAAA	960
	TTTTTCAATT					300
AGTGATTCTA	GCAATAGTGA	TTCCACTCAG	AGTCAGAAGT	CTGGCAGGAA	CTCGAACCCC	1020
TCACTAAGAT	CGTTATCACT	AAGGTGAGTC	TCAGTCTTCA	GACCGTCCTT	GAGCTTGGGG	
	GCAACTAAAT					1080
GCCGTTCGTG	CGTTGATTTA	GGGCTTTATG	TTTTTCATTG	TGTCACCTGA	AGGATAATTC	
ACTTACTTGC	ATTGCTGGAC	TAGCAAAGGA	AAATTGCACT	ልጥጥርርልሮልጥር	አ ጥአጥጥርጥአጥጥ	1140
	TAACGACCTG					1140
GTTTACTATA	AAAATCATGT	GATAACTGAT	TATTACTTCT	GTTTCTCTTT	TGGTTTCTGC	1200
CAAATGATAT	TTTTAGTACA	CTATTGACTA	ATAATGAAGA	CAAAGAGAAA	ACCAAAGACG	
TTCTCTCTTC	TCTCAACCCC	ጥጥጥርጥል ልጥርር	ጥጥርር ርርርር ል	CACTCTTAAC	ጥ አ ጥአ ጥጥር ጥር አ	1260
	AGAGTTGGGG					1200
	manamana	maranana	mammammma.			4000
	TCACTAATCA AGTGATTAGT					1320
CHRIMONIAN	NGIGNIINGI	ACICITIIG	ACAAGAAAAC	GITATTATTA	TTTAATTTGT	
	AGAGCCTCTT					1380
ACGACAATGG	TCTCGGAGAA	ACGACTCAGA	GGTCTACAAT	TAAATGAAAG	ACGTGGGGTT	
TTGGGAATGC	AATATTGGAT	GAAAAGAGAG	GTTTCTGGTA	TTCACAGAAA	GCTAGATATG	1440
AACCCTTACG	TTATAACCTA	CTTTTCTCTC	CAAAGACCAT	AAGTGTCTTT	CGATCTATAC	
ССТТААААСА	TACTCTGCCG	ΑͲϹͲΑΑͲͲΑϹ	ልርረርጥጥልጥጥጥ	ጥጥሩ ጥልጥሩርርጥ	ጥጥ ጉርረርር ለ ውጥ	1500
	ATGAGACGGC					1500
CTCCTCATGC	TTAGAAAGTT	CCAAATGTTT	ATAAAGGTAA	AATGGCAGTT	TGAAGTCAAA	1560
	AATCTTTCAA					2500
					ACACCCAAGA	1620
ACAGTGTATC	CGTTTCGTTA	GTTCGTGGTC	CTTCACAAAT	ACTCCTTTGT	TGTGGGTTCT	
TGAATTATTT	TTGAGACTGT	CAGGAAGTAA	AATAAATAGG	AGCTTAAGAA	AGAACATTTT	1680
					TCTTGTAAAA	
GCCTGATTGA	GAAGCACAAC	TGAAACCAGT	AGCCGCTGGG	GTGTTAATGG	TAGCATTCTT	1740
CGGACTAACT	CTTCGTGTTG	ACTTTGGTCA	TCGGCGACCC	CACAATTACC	ATCGTAAGAA	
CTTTTGGC 2	ጥ ልርልጥጥጥርኔጥ	<u> ጥጥርምጥር ልጥር ል</u>	ል ጥልጥልጥጥል ልጥ	<u>ሮ</u> ልርር አጥጥ አር አ	GAAATGAATT	1800
					CTTTACTTAA	1000
						د
					TTTAAATAAA	1860
TATIGATCIG	TAGACGACAA	TAGTGGTATC	AAAACAAATT	AAACGAAGGA	AAATTTATTT	
CCCATTGGTG	AAAGTCAAAA	АААААААА	AAA			
GGGTAACCAC	TTTCAGTTTT	TTTTTTTTT	TTT			

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